

Table S2: Information on the *Anopheles* clusters analysed in this work

miRNA Name	miRNA coordinates			Strand	Distance (kb)	Hosting protein coding gene	Predicted ^a
	Chromosome	Start	End				
miR-281*	2L	17362447	17362467	+	0		W, M
miR-J*	2R	9129644	9129665	-	0		
miR-G*	2R	9129834	9129857	-	0.17		
miR-34*	2R	28232721	28232740	-	0		W
miR-277	2R	28234532	28234554	-	1.8		W, M
miR-317	2R	28252007	28252031	-	20.2		W, M
miR-12*	2R	37888125	37888147	+	0	<i>ENSANGG00000011107</i>	
miR-283	2R	37890092	37890112	+	2	<i>ENSANGG00000011107</i>	W, M
miR-F*	2R	55572844	55572865	-	0	<i>ENSANGG00000026438</i>	
miR-279	2R	55573339	55573360	-	0.5	<i>ENSANGG00000026438</i>	W, M
miR-I*	3L	2905485	2905505	-	0		
miR-306*	3R	5888646	5888667	-	0	<i>ENSANGG00000008044</i>	
miR-79	3R	5888312	5888333	-	0.3	<i>ENSANGG00000008044</i>	W, M
miR-9b	3R	5887842	5887864	-	0.7	<i>ENSANGG00000008044</i>	W, M
miR-9c	3R	5891223	5891244	-	7.5	<i>ENSANGG00000008044</i>	W, M
let-7*	3R	10270762	10270783	-	0		W, M
miR-125	3R	10270221	10270242	-	0.5		W, M
miR-100	3R	10274733	10274754	-	3.9		W, M
miR-275*	3R	38377402	38377423	+	0		W, M
miR-305	3R	38386260	38386282	+	8.8		W, M

* miRNA cloned in this study

^a The miRNA has been predicted by Wang et al., 2005 (W) or in the miRBase (M)